

GenCore version 4.5
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OM protein - protein search, using sw model

Run on: August 28, 2002, 17:32:27 ; Search time 37.53 seconds

(without alignments)
637.523 Million cell updates/sec

Title: US-09-502-984B-37

Perfect score: 1284

Sequence: 1 KESKKAALLAARGPEELLCF.....RKMERLEEEVERLKLQVGER 249

Scoring table:

BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 283138 seqs, 96089334 residues

Total number of hits satisfying chosen parameters: 283138

Minimum DB seq length: 0
Maximum DB seq length: 2000000000Post-processing: Minimum Match 0%
Maximum Match 100%

Listing first 45 summaries

Database :

1: pir1.*
2: pir2.*
3: pir3.*
4: pir4.*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Match	Query Length	DB ID	Description
1	1060	82.6	508	1 ZUHUR	erythropoietin rec
2	877.5	68.3	507	2 A46713	erythropoietin rec
3	875.5	68.2	265	2 S14081	erythropoietin rec
4	869.5	67.7	507	1 A32385	erythropoietin rec
5	171	13.3	625	2 S35317	hematopoietic grow
6	164	12.8	626	2 S37622	proto-oncogene - m
7	146	11.4	579	2 B45266	MPL-K protein prec
8	146	11.4	635	2 A45266	MPL-K protein prec
9	118	9.2	581	2 I45971	interleukin-9 rece
10	114.5	8.9	522	2 B45268	interleukin-9 rece
11	110.5	8.6	634	2 S33339	somatotropin recep
12	110	8.6	831	2 J01655	somatotropin recep
13	108	8.4	281	1 RGBY22	prolactin receptor
14	107	8.3	608	2 S32822	amino acid biosynt
15	104	8.1	616	2 A30304	somatotropin recep
16	103.5	8.1	677	2 S33608	prolactin receptor
17	103.5	8.1	830	2 I50455	somatotropin-bindi
18	103	8.0	638	2 S12136	prolactin receptor
19	99.5	7.7	638	2 B28176	somatotropin recep
20	98	7.6	897	1 A39255	cytokine receptor
21	96.5	7.5	378	2 A40266	interleukin-3 rece
22	95	7.4	814	2 A62051	endopeptidase C1p
23	94	7.3	468	2 A45268	interleukin-9 rece
24	94	7.3	1825	2 C88400	protein H19M22.1 f
25	94	7.3	1825	2 T32828	hypothetical prote
26	92.5	7.2	279	2 B32985	somatotropin-bindi
27	92.5	7.2	638	2 A33505	somatotropin recep
28	92	7.2	2594	2 A57774	kinase-related pro
29	91	7.1	284	2 S27931	Env/v-mpl fusion p

30	91	7.1	292	2 I77525	prolactin receptor
31	91	7.1	303	2 I77524	prolactin receptor
32	91	7.1	608	2 I53269	prolactin receptor
33	90	7.0	467	2 I56896	gene gfi-2 protein
34	90	7.0	485	1 K1BYHA	hexokinase (EC 2.7
35	90	7.0	682	1 H69879	ATP-dependent DNA
36	88.5	6.9	511	2 A37803	beta-fructofuranos
37	88	6.9	288	2 B59405	prolactin receptor
38	88	6.9	376	2 A59405	prolactin receptor
39	88	6.9	622	2 A40144	prolactin receptor
40	88	6.9	841	2 A43254	protein-tyrosine-p
41	87	6.8	310	2 A29884	prolactin receptor
42	87	6.8	412	2 A41070	prolactin receptor
43	87	6.8	610	2 A34631	lactogen receptor
44	87	6.8	610	2 A36116	prolactin receptor
45	86.5	6.7	406	2 F70019	nifs protein homol

ALIGNMENTS

RESULT 1

ZUHUR

erythropoietin receptor precursor - human

C:Species: Homo sapiens (man)

C:Date: 12-Feb-1993 #sequence_revision 05-Apr-1995 #text_change 22-Jun-1999

C:Accession: A43799; A60160; A49824; A53958; A55280; I52563

R:Jones, S.S.; D'Andrea, A.D.; Haines, L.L.; Wong, G.G.

Blood 76, 31-35, 1990

A:Title: Human erythropoietin receptor: cloning, expression, and biologic characteriz

A:Reference number: A43799; MUID:90304340

A:Accession: A43799

A:Molecule type: mRNA

A:Residues: 1-508 <CON>

A:Cross-references: GB:M60459; NID:q182244; PIDN:AAA52403.1; PID:q182245

R:Winkelmann, J.C.; Penny, L.A.; Deaven, L.L.; Forget, B.G.; Jenkins, R.B.

Blood 76, 24-30, 1990

A:Title: The gene for the human erythropoietin receptor: analysis of the coding seque

A:Reference number: A60160; MUID:90304334

A:Accession: A60160

A:Status: not compared with conceptual translation

A:Molecule type: mRNA; DNA

A:Residues: 1-101, 'R', 103-188, 'RP', 191-243, 'E', 245-508 <MIN>

R:Noguchi, C.T.; Bae, K.S.; Chiu, K.; Wade, Y.; Schechter, A.N.; Hankins, W.D.

Blood 78, 2548-2556, 1991

A:Title: Cloning of the human erythropoietin receptor gene.

A:Reference number: A49824; MUID:92399733

A:Accession: A49824

A:Molecule type: DNA

A:Residues: 1-508 <NOG>

A:Cross-references: GB:S45332; NID:q255496; PIDN:AA23271.1; PID:q255497

A:Experimental source: Placenta

A>Note: sequence extracted from NCBI backbone (NCBIN:113293, NCBIPI:113294)

R:Ehrenman, K.; St. John, T.

Exp. Hematol. 19, 973-977, 1991

A:Title: The erythropoietin receptor gene: cloning and identification of multiple tra

A:Reference number: A53958; MUID:91372559

A:Accession: A53958

A:Molecule type: mRNA

A:Residues: 1-508 <EHR>

R:Penny, L.A.; Forget, B.G.

Genomics 11, 974-980, 1991

A:Title: Genomic organization of the human erythropoietin receptor gene.

A:Reference number: A55280; MUID:92147143

A:Accession: A55280

A:Molecule type: DNA

A:Residues: 1-17,381-387, 'LLEQQQDA', 391-395,504-508 <PEN>

A>Note: sequence modified after extraction from NCBI backbone

A>Note: an insert compared to other published sequences is considered by authors as 1

R:Maouche, L.; Tournamille, C.; Hattab, C.; Boffa, G.; Cartlon, J.P.; Chretien, S.

Blood 78, 2557-2563, 1991

A:Title: Cloning of the gene encoding the human erythropoietin receptor.

A:Reference number: 152563; MUID:92399734
 A:Accession: 152563
 A:Status: translated from GB/EMBL/DBJ
 A:Molecule type: DNA
 A:Residues: 1-96 <RES>
 A:Cross-references: GB:M76595; NID:q182147; PIDN:AAA52393.1; PID:q553281
 C:Genetics:
 A:Gene: GDB:EPOR
 A:Cross-references: GDB:125242; OMIM:133171
 A:Map position: 19p13.3-19p13.2
 A:Introns: 33/1; 84/2; 143/1; 195/3; 247/1; 276/2; 305/3
 C:Superfamily: erythropoietin receptor; cytokine receptor; glycoprotein; transmembrane protein
 C:Keywords: alternative splicing; cytokine receptor; glycoprotein; transmembrane protein
 F:1-24/Domain: signal sequence #status predicted <SIG>
 F:25-508/Product: erythropoietin receptor #status predicted <EXT>
 F:52-250/Domain: extracellular #status predicted <EXT>
 F:52-239/Domain: cytokine receptor homology <CRS>
 F:233-237/Region: WSXWS motif
 F:251-272/Domain: transmembrane #status predicted <TM>
 F:273-508/Domain: intracellular #status predicted <INT>
 F:52-62,91-107/Disulfide bonds: #status predicted
 F:76/Binding site: carbohydrate (Asn) (covalent) #status predicted

Query Match 82.6%; Score 1060; DB 1; Length 508;
 Best Local Similarity 93.8%; Pred. No. 2,4e-84;
 Matches 198; Conservative 11; Mismatches 2; Indels 0; Gaps 0;

QY 1 KFSKALLAARPEELCTERLEEDVCFEEAASAGVPGNFSFOLEDEPMKRL 60
 |||||
 Db 34 KFSKALLAARPEELCTERLEEDVCFEEAASAGVPGNFSFOLEDEPMKRL 93
 QY 61 HQAPTARGAIRFWCSLPTADTSSFPVLELRLTAASGAPRFRVHIINEVLLDAPVGLVA 120
 |||||
 Db 94 HQAPTARGAIRFWCSLPTADTSSFPVLELRLTAASGAPRFRVHIINEVLLDAPVGLVA 153
 QY 121 RLADSGHVYIRLPPPEPMTSHIRELDISANGAGSVQVRELLGRTCYLSNLGR 180
 |||||
 Db 154 RLADSGHVYIRLPPPEPMTSHIRELDISANGAGSVQVRELLGRTCYLSNLGR 213
 QY 181 TRTIVARMAEPSPFGFMSANSEPSLTLT 211
 |||||
 Db 214 TRTIVARMAEPSPFGFMSANSEPSLTLT 244

RESULT 2
 A46713
 erythropoietin receptor precursor - rat
 C:Species: Rattus norvegicus (Norway rat)
 C>Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 16-Jun-2000
 C:Accession: A46713
 R:Masuda, S.; Nagao, M.; Takahata, K.; Konishi, Y.; Gallyas Jr., F.; Tabira, T.; Sakaki, J. Biol. Chem. 268, 11208-11216, 1993
 A:Title: Functional erythropoietin receptor of the cells with neural characteristics. C
 A:Reference number: A46713; MUID:9326574
 A:Accession: A46713
 A:Status: preliminary
 A:Molecule type: mRNA
 A:Residues: 1-507 <MAS>
 A:Cross-references: GB:D13566; NID:q286209; PIDN:BA02761.1; PID:q286210
 A:Experimental source: PC12 and erythroid cells
 A:Note: sequence extracted from NCBI backbone (NCBI:132811, NCBI:132813)
 C:Superfamily: erythropoietin receptor; cytokine receptor homology
 C:Keywords: cytokine receptor; glycoprotein; transmembrane protein
 F:1-24/Domain: signal sequence #status predicted <SIG>
 F:25-507/Product: erythropoietin receptor #status predicted <MAT>
 F:25-249/Domain: extracellular #status predicted <EXT>
 F:52-238/Domain: cytokine receptor homology <CRS>
 F:250-271/Domain: transmembrane #status predicted <TM>
 F:272-507/Domain: intracellular #status predicted <INT>
 F:75/Binding site: carbohydrate (Asn) (covalent) #status predicted

Query Match 68.3%; Score 877.5; DB 1; Length 507;
 Best Local Similarity 78.2%; Pred. No. 1.8e-68;
 Matches 165; Conservative 23; Mismatches 22; Indels 1; Gaps 1;

QY 1 KFSKALLAARPEELCTERLEEDVCFEEAASAGVPGNFSFOLEDEPMKRL 60
 |||||
 Db 34 KFSKALLAARPEELCTERLEEDVCFEEAASAGVPGNFSFOLEDEPMKRL 92
 QY 61 HQAPTARGAIRFWCSLPTADTSSFPVLELRLTAASGAPRFRVHIINEVLLDAPVGLVA 120
 |||||
 Db 93 HQAPTARGAIRFWCSLPTADTSSFPVLELRLTAASGAPRFRVHIINEVLLDAPVGLVA 152
 QY 121 RLADSGHVYIRLPPPEPMTSHIRELDISANGAGSVQVRELLGRTCYLSNLGR 180
 |||||
 Db 153 RLADSGHVYIRLPPPEPMTSHIRELDISANGAGSVQVRELLGRTCYLSNLGR 212
 QY 181 TRTIVARMAEPSPFGFMSANSEPSLTLT 211
 |||||
 Db 213 TRTIVARMAEPSPFGFMSANSEPSLTLT 243

RESULT 3
 S14081

erythropoietin receptor - mouse
 C:Species: Mus musculus (house mouse)
 C>Date: 18-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
 C:Accession: S14081; I49653
 R:Kuramochi, S.; Ikawa, Y.; Todokoro, K.
 J. Mol. Biol. 216, 567-575, 1990
 A:Title: Characterization of murine erythropoietin receptor genes.
 A:Reference number: S13249; MUID:91080149
 A:Accession: S14081

A:Status: preliminary
 A:Molecule type: DNA
 A:Residues: 1-265 <KUR>
 R:Jacobs, C.; Chretien, S.; Lemarchandel, V.; Mayeux, P.; Romeo, P.
 J. Biol. Chem. 266, 6952-6956, 1991
 A:Title: Spleen focus-forming virus long terminal repeat insertional activation of th
 A:Reference number: I49653; MUID:91201346
 A:Accession: I49653

A:Status: preliminary; translated from GB/EMBL/DBJ
 A:Molecule type: DNA
 A:Residues: 1-24 <RES>
 A:Cross-references: GB:M62360; NID:q193199; PIDN:AAA37582.1; PID:q193200
 C:Superfamily: erythropoietin receptor; cytokine receptor homology
 C:Keywords: cytokine receptor; transmembrane protein
 F:52-238/Domain: cytokine receptor homology <CRS>

Query Match 68.2%; Score 875.5; DB 2; Length 265;
 Best Local Similarity 77.1%; Pred. No. 1.2e-68;
 Matches 165; Conservative 22; Mismatches 26; Indels 1; Gaps 1;

QY 1 KFSKALLAARPEELCTERLEEDVCFEEAASAGVPGNFSFOLEDEPMKRL 60
 |||||
 Db 34 KFSKALLAARPEELCTERLEEDVCFEEAASAGVPGNFSFOLEDEPMKRL 92
 QY 61 HQAPTARGAIRFWCSLPTADTSSFPVLELRLTAASGAPRFRVHIINEVLLDAPVGLVA 120
 |||||
 Db 93 HQAPTARGAIRFWCSLPTADTSSFPVLELRLTAASGAPRFRVHIINEVLLDAPVGLVA 152
 QY 121 RLADSGHVYIRLPPPEPMTSHIRELDISANGAGSVQVRELLGRTCYLSNLGR 180
 |||||
 Db 153 RLADSGHVYIRLPPPEPMTSHIRELDISANGAGSVQVRELLGRTCYLSNLGR 212
 QY 181 TRTIVARMAEPSPFGFMSANSEPSLTLTGGG 214
 |||||
 Db 213 TRTIVARMAEPSPFGFMSANSEPSLTLTASG 246

RESULT 4
 A32385
 erythropoietin receptor precursor, membrane-bound form - mouse

C.Species: Mus musculus (house mouse)
C.Date: 28-Sep-1990 #sequence_revision 05-Apr-1995 #text_change 22-Jun-1999
C.Accession: A41686; A32385; S13249
R.Hino, M.; Tojo, A.; Misawa, Y.; Morii, H.; Takaku, F.; Shibuya, M.
Mol. Cell. Biol. 11, 5527-5533, 1991
A.Title: Unregulated expression of the erythropoietin receptor gene caused by insertion
A.Reference number: A41686; MUID:92017832
A.Accession: A41686
A.Molecule type: mRNA
A.Residues: 1-507 <HIN>
A.Cross-references: GB:S59388; NID:g237036; PIDN:AAB20029.1; PID:g237037
A.Experimental source: murine erythroleukemia (MEL) cell line F5-5
R.D'Andrea, A.D.; Lodish, H.F.; Wong, G.G.
Cell 57, 277-285, 1989
A>Title: Expression cloning of the murine erythropoietin receptor.
A.Reference number: A32385; MUID:89195238
A.Accession: A32385
A.Molecule type: mRNA
A.Residues: 1-507 <DNA>
A.Cross-references: GB:J04843; NID:g193090; PIDN:AAA37571.1; PID:g309219
A.Experimental source: murine erythroleukemia (MEL) cells, subclone 745
R.Kuramotochi, S.; Ikawa, Y.; Todokoro, K.
J. Mol. Biol. 216, 567-575, 1990
A>Title: Characterization of murine erythropoietin receptor genes.
A.Reference number: S13249; MUID:91080149
A.Accession: S13249
A.Molecule type: DNA; mRNA
A.Cross-references: EMBL:X51081; NID:g50861; PIDN:CA37248.1; PID:g50862
A.Experimental source: murine erythroleukemia K-1 cells
C.Genetics:
A.Introns: 39/1; 83/2; 142/1; 194/3; 246/1; 275/2; 304/3
C.Superfamily: erythropoietin receptor; cytokine receptor homology
C.Keywords: alternative splicing; cytokine receptor; glycoprotein; transmembrane protein
F.1-24/DNA: signal sequence #status predicted <SIG>
F.25-507/Product: erythropoietin receptor #status predicted <MAT>
F.25-248/DNA: extracellular #status predicted <EXT>
F.250-238/DNA: cytokine receptor homology <CRS>
F.250-238/DNA: transmembrane #status predicted <TM>
F.272-507/DNA: intracellular #status predicted <INT>
F.52-62, 90-106/Disulfide bonds: #status predicted
F.75/Binding site: carbohydrate (Asn) (covalent) #status predicted

Query Match 67.7%; Score 869.5; DB 1; Length 507;
Best Local Similarity 77.7%; Pred. No. 9e-68;
Matches 164; Conservative 22; Mismatches 24; Indels 1; Gaps 1;

OY 1 KFESAAALLAARGPELLCTFERLEDVCFEEAASAGVGPGNFSEFOLEDEPWKLCRL 60
 ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 34 KFESAAALLAARGSELLCTFORLEDVCFWEAASSGM-DFNYSFYQLEGSRKSSCL 92
 ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | |
OY 61 HOAPARAARPCWGLPADRSSFPVLELRILTAAGCARFRHVTHINEVILLDPAVGVA 120
 ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 93 HOAPVRKSVKWCGLPADRSSFVPLELYQTVEASGSPRYHRIITHINEVILLDAPAGLA 152
 ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | |
OY 121 RLADSGHHVIRFWLPPEPTMTSHIRFELDLSAGAGSAVSORVELLEERTCECVSNLNGR 180
 ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 153 RRAEGSHIVLRMLPPPGAPMTHTIRYEVDVASGRAGACTOREVLEBRTCECVSNLNGG 212
 ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | |
OY 181 TRITIAVARMAEPSEFGFSAMSEPSVLTT 211,
 || | | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 213 TRYTPAVBARMAEPSFGFSAMSEPSVLTT 243
 || | | | | | | | | | | | | | | | | | | | | | | | | | | | |

RESULT 5
S35317
hematopoietic growth factor receptor precursor - mouse
C.Species: Mus musculus (house mouse)
C.Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 05-Nov-1999
C.Accession: S35317; S35316
C.Rambo, J.; Selvidj, D.C.; Chiang, M.K.; Feichel, C.L.; Vogt, T.F.; Leder, P.
EMBO J. 12, 2645-2653, 1993

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A>Title: Murine c-mpl: a member of the hematopoietic growth factor receptor superfamily
A:Reference number: S35316; MUID:93327753
A:Accession: S35317
A:Molecule type: mRNA
A:Residues: 1-625 <SKO>
A:Cross-references: EMBL:Z22649; NID:g394725; PIDN:CAAB0365.1; PID:g394726
A:Accession: S35316
A:Molecule type: DNA
A:Residues: 1-70 <SKM>
A:Cross-references: EMBL:Z22657
C:Keywords: alternative splicing; cytokine receptor; glycoprotein; transmembrane prot
F:1-25/Domain: signal sequence #status predicted <SIG>
F:26-625/Product: hematopoietic growth factor receptor #status predicted <MAT>
F:26-482/Domain: extracellular #status predicted <EXT>
F:261-265/Region: WGXWS motif
F:465-469/Region: WSXWS motif
F:483-504/Domain: transmembrane #status predicted <TMW>
F:505-625/Domain: intracellular #status predicted <INT>
F:113,117,178,349/Binding site: carbohydrate (asn) (covalent) #status predicted

Query Match      13.3%; Score 171; DB 2; Length 625;
Best Local Similarity 24.6%; Pred. No. 6.4e-07;
Matches 62; Conservative 33; Mismatches 105; Indels 52; Gaps 8;

QY 4 SKAALLARGPPELLCTFERLEDLVCFPEEASAGVGNGNSFSFOLEDEPMKLCRLHOA 63
   | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
DB 25 SODVFLALGTRPLNCFCOTFEDLTGCFWDEBEAA--PGTGQLLXATRGKPRACPLYSQ 82

QY 64 PARGAIRFWCSLPAD-TTSFVPLELRITAA-SGARFHRYIHINEVYLDAVPGLVAR 121
   | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
DB 83 SVPTGRTGRVCOFPADDEVRLFELHWNKNVSLNOTLIQHVLVDSDGLPAPRPVIKAR 142

QY 122 LADSGHVYIR-LRPPETPMHSIRELDI-----SAGNGAGSYQRVELLEGRI----- 170
   | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
DB 143 GGSQPCQLDIHEADAPE--ISDFLRHLRYGPFDDSSNAATPSV--IQLLSTETCCPTLW 198

QY 171 -----ECVLSNLGGRRITRTAVPARMAEPSFG 197
   | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
DB 199 MNPNPVVLDOQPCPHPTASQPHGRAPFLTVGGSCLVSGDASNSYNWLQRSQDGVSILR 258

QY 198 GFWSAMSEPVSL 209
   | | | | | : | : | : | : | : | : | : | : | : | : | : | : | : |
DB 259 GSMGPMSEPVTV 270

RESULT 6
S37622
proto-oncogene - mouse
C.Species: Mus musculus (house mouse)
C.Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 05-Nov-1999
C.Accession: S37622
R.Vignon, I.; Florindo, C.; Fichelson, S.; Guenet, J.L.; Mattei, M.G.; Souyri, M.; Cos
Onogene 8, 2607-2615, 1993
A.Title: Characterization of the murine Mpl proto-oncogene, a member of the hematoipo
A:Reference number: S37622; MUID:93390934
A:Accession: S37622
A>Status: Preliminary
A:Molecule type: mRNA
A:Residues: 1-626 <VIG>
A:Cross-references: EMBL:X73677; NID:g404318; PIDN:CAM52031.1; PID:g404319

Query Match      12.8%; Score 164; DB 2; Length 626;
Best Local Similarity 23.5%; Pred. No. 2.6e-06;
Matches 61; Conservative 34; Mismatches 105; Indels 60; Gaps 8;

QY 4 SKAALLARGPPELLCTFERLEDLVCFPEEASAGVGNGNSFSFOLEDEPMKLCRLHOA 63
   | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
DB 18 SODVFLALGTRPLNCFCOTFEDLTGCFWDEBEAA--PSGTQLLYAYARGEKPRACPLYSQ 75

QY 64 PARGAIRFWCSLPAD-TTSFVPLELRITAA-SGARFHRYIHINEVYLDAVPGLVAR 121

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[illegible]

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Query Match 59: 11.4%; Score 146; DB 2; Length 579;
Best Local Similarity 22.8%; Pred. No. 8.8e-05;
Matches 59; Conservative 35; Mismatches 107; Indels 58; Gaps 9;

QY 4 SKAALLAARGPPELLCTFERLEDLVCFEEENASAGVGGNGFSFSQLEDEPKLCRLHQA 63
      | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
DB 25 SODVSLASDSEPLKCFSTFEDLTCFWDDEEAAASGYQLLYAIPRE-KP-RACPSSSQ 82
      | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
QY 64 PTARCAIFWCSLP-TADTSSFVPLELR-LTAASGAPRHRVYTHINEVLLDAPYGLVAR 121
      | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
DB 83 SMPHGETGYVCOFPDQEEVRLFFEPFLHWKNVFLNQTRQRYLFEVDSGLPAPPSIIKAM 142
      | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
QY 122 LADESGHVIRN-LRPPEPMTHIRFEL-----LISAGNGASVQRE----- 164
      | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
DB 143 GGSQGGELQISWEEPAPE-ISRDLRYELRYGPRDPKNSGTGYQLLATETCCPALQRP 200
      | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
QY 165 -----LLEGTECVLSNLGORTITTAIVRAR 190
      | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
DB 201 HSASALDPSPCAQPIPMQDQPKQTSFRESALTAIEGGS-CLISGLQPGNSYWLQLNSE 259
      | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
QY 191 MAEPFSGGFWSAMSEPSVL 209
      | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
DB 260 PDGISLGGSMGSMSLPPTY 278
      | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |

RESULT 8
A45266
MPL-P protein precursor - human
C:Species: Homo sapiens (man)
C:Date: 27-Jun-1994 #sequence_revision 27-Jun-1994 #text_change 05-Nov-1999
C:Accession: A45266
R:Vignou, I.; Morron, J.P.; Coccault, L.; Mitjavila, M.T.; Tambourin, P.; Gisselbrecht, S
Proc. Natl. Acad. Sci. U.S.A. 89, 5640-5644, 1992
A:Title: Molecular cloning and characterization of MPL, the human homolog of the v-mpl c
A:Reference number: A45266; MUID:92302297
A:Accession: A45266
A:Status: Preliminary
A:Molecule type: mRNA
A:Residues: 1-635 <VIG>
A:Cross-references: GB:M90102; NID:g184260; PIDN:AAA69971.1; PID:g184261

```

C;Keywords: transmembrane protein

```
Query Match      11.4%; Score 146; DB 2; Length 635;  
Best Local Similarity    22.8%; Pred. No. 9,8e+05;  
Matches     59; Conservative   35; Mismatches 107; Indels   58; Gaps  
  
QY       4 SKAALLARGPELLCTFETRLLEDLVCFEEENASAGVGPNGFSFOLEDEPWKLCRLHQ 63  
          | : : : : : : : : : : : : : : : : : : : : : : : :  
Db       25 SQDVSLIASDSBP LKCFERTFEDLTCFMDEEAAPSGTQLLYAVPRE-KP-RACPISSQ 82  
  
QY       64 PTARGAIRFWCSLP-TADTSSFPVLRL-TAACAPRPHRYIHNVEVLDPAYGLVAR 121  
          | : : : : : : : : : : : : : : : : : : : : : : : :  
Db       83 SMPHGRTYVCQCPDQEVRLEFFPLHWKNVFNLQTGFVLFVDVGALPAPSIIKAM 142  
  
QY       122 LADEGHHVIYIK-LPPETPMTSIRTEL-----DISAGNAGSVQRVE----- 164  
          | : : : : : : : : : : : : : : : : : : : : : : : :  
Db       143 GGSQCPELQISWEEPAPE--ISDFRLRYELRGGRPKNKSTGGTVIQLLATETCCPALORP 200  
  
QY       165 -----LLEGRTGVLSNLNRGTRITIAVRAR 190  
          || : : : : : : : : : : : : : : : : : : : : : : : :  
Db       201 HSASALDOSPCAQPTMPWDQGPKTSPSRASAALTAEGBS-CLISGLQPGNSYIMQLNSE 259  
  
QY       191 MAEPFGGFMSAMSSEPVSL 209  
          | | | | : : | : : : :  
Db       260 PDGISLGSGWSGWSMLPVTIV 278
```

RESULT 9
145971
prolactin receptor - bovine
C:Species: Bos primigenius taurus (cattle)
C:Date: 16-Aug-1996 #sequence_revision 16-Aug-1996 #text_change 28-Jul-2000
C:Accession: 145971
R:Scott, P.; Kessler, M.A.; Schuler, L.A.
Mol. Cell. Endocrinol. 89, 47-58, 1992
A1:Title: Molecular cloning of the bovine prolactin receptor and distribution of prolactin receptor mRNA
A:Reference number: 145971; MUID:93246019
A:Accession: 145971
A:Status: preliminary; translated from GB/EMBL/DBD
A:Molecule type: mRNA
A:Residues: 1-581 <SC0>
A:Cross-references: GB:I02549; NID:g163617; PIDN:AAA51417.1; PID:g163618
C:Genetics:
A:Gene: PRLR
C:Superfamily: cytokine receptor homology
i:36-221/Domain: cytokine receptor homology <CRS>

Query Match	21.9%	Score 118;	DB 2;	Length 581;
Best Local Similarity	21.9%	Pred. No. 0.024;		
Matches	48;	Conservative	40;	Mismatches 101; Indels 30; Gaps 10;

QY	7	ALLAARGEE---	LLCFETLERLEDVCFEEAASAGVGGNFSFSFOLEDEPWKLCRLHQ	62
DB	20	SILNQSPPEKFKLVKCRSPGKEFTFCWMEPCADGGL-PTNLTLYHKKGE---	TLIHE	74
QY	63	APTARGAIRFMOCLPTADSSFPVLELRFTA-----	ASGAPRFRHVIHINEVLLDADAV	116
DB	75	CPDYTGCGPNCSTFESKRTSTIKMYVIITYNAINOMGSISSDPL---	VHVTIIVEEPFA	131
QY	117	GLVARL---ADESGHVIVIRKLPPETPMTSH--	IRFELDISAGNAGSGSVORVELLEGRT	170
DB	132	NLTLELKHEDRKPRLMIKWSPTMTDVSQGAFFIIQYELRLPERKATDMETHFLTKO--T	189	
QY	171	ECVLSNLGRFRPLITAVARRMAEPRSGGWSMSAEVSL	209	
DB	130	QKTIENLTPGQKYLVAQIRCK--PDH-GIWSKMSPPSSI	224	

RESULT 10
B45268
interleukin-9 receptor precursor - human

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